



SEQUENCE LISTING

<110> Clarkson, Kathleen A.
Fenel, Fred

<120> Modified Enzymes, Methods to Produce
Modified Enzymes and Uses Thereof

<130> GC812-C1

<140> US 11/404,460

<141> 2006-04-14

<150> US 10/565,954

<151> 2004-09-10

<150> US 60/503,251

<151> 2003-09-15

<160> 51

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 223

<212> PRT

<213> Trichoderma reesei

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Arg	Gln	Thr	Ile	Gln	Pro	Gly	Thr	Gly	Tyr	Asn	Asn	Gly	Tyr	Phe	Tyr
		35					40					45			
Ser	Tyr	Trp	Asn	Asp	Gly	His	Gly	Gly	Val	Thr	Tyr	Thr	Asn	Gly	Pro
	50					55					60				
Gly	Gly	Gln	Phe	Ser	Val	Asn	Trp	Ser	Asn	Ser	Gly	Asn	Phe	Val	Gly
65						70				75				80	
Gly	Lys	Gly	Trp	Gln	Pro	Gly	Thr	Lys	Asn	Lys	Val	Ile	Asn	Phe	Ser
				85					90					95	
Gly	Ser	Tyr	Asn	Pro	Asn	Gly	Asn	Ser	Tyr	Leu	Ser	Val	Tyr	Gly	Trp
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Ser	Arg	Asn	Pro	Leu	Ile	Glu	Tyr	Tyr	Ile	Val	Glu	Asn	Phe	Gly	Thr
		115					120					125			
Tyr	Asn	Pro	Ser	Thr	Gly	Ala	Thr	Lys	Leu	Gly	Glu	Val	Thr	Ser	Asp
	130					135					140				
Gly	Ser	Val	Tyr	Asp	Ile	Tyr	Arg	Thr	Gln	Arg	Val	Asn	Gln	Pro	Ser
145					150					155				160	
Ile	Ile	Gly	Thr	Ala	Thr	Phe	Tyr	Gln	Tyr	Trp	Ser	Val	Arg	Arg	Asn
				165					170					175	
His	Arg	Ser	Ser	Gly	Ser	Val	Asn	Thr	Ala	Asn	His	Phe	Asn	Ala	Trp
			180					185					190		
Ala	Gln	Gln	Gly	Leu	Thr	Leu	Gly	Thr	Met	Asp	Tyr	Gln	Ile	Val	Ala
		195					200						205		

Val Glu Gly Tyr Phe Ser Ser Gly Ser Ala Ser Ile Thr Val Ser
 210 215 220

<210> 2
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 <212> DNA
 <213> Trichoderma reesei

<400> 2
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 ggctacaaca acggctactt ctactcgtae tggaacgatg gccacggcgg cgtgacgtac 180
 accaatgggc cggcggggca gttctccgtc aactgggtcca actcggggcaa ctttgtcggc 240
 ggcaagggat ggcagcccgg caccaagaac aagtaagact acctactctt accccctttg 300
 accaacacag cacaacacaa tacaacacat gtgactacca atcatggaat cggatctaac 360
 agctgtgttt tcaaaaaaaaaa gggatcatcaa cttctcgggc agctacaacc ccaacggcaa 420
 cagctacctc tccgtgtacg gctgggtccc caacccccctg atcgagtact acatcgtcga 480
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 cggcagcgtc tacgacattt accgcacgca gcgcgtcaac cagccgtcca tcatcggcac 600
 cgccaccttt taccagtact ggtccgtccg ccgcaaccac cgctcgagcg gctccgtcaa 660
 cagggcgaac cacttcaacg cgtggggtca gcaaggcctg acgctcggga cgatggatta 720
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<210> 3
 <211> 234
 <212> PRT
 <213> Trichoderma reesei

<400> 3
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 Gln Thr Ser Cys Asp Gln Trp Ala Thr Phe Thr Gly Asn Gly Tyr Thr
 20 25 30
 Val Ser Asn Asn Leu Trp Gly Ala Ser Ala Gly Ser Gly Phe Gly Cys
 35 40 45
 Val Thr Ala Val Ser Leu Ser Gly Gly Ala Ser Trp His Ala Asp Trp
 50 55 60
 Gln Trp Ser Gly Gly Gln Asn Asn Val Lys Ser Tyr Gln Asn Ser Gln
 65 70 75 80
 Ile Ala Ile Pro Gln Lys Arg Thr Val Asn Ser Ile Ser Ser Met Pro
 85 90 95
 Thr Thr Ala Ser Trp Ser Tyr Ser Gly Ser Asn Ile Arg Ala Asn Val
 100 105 110
 Ala Tyr Asp Leu Phe Thr Ala Ala Asn Pro Asn His Val Thr Tyr Ser
 115 120 125
 Gly Asp Tyr Glu Leu Met Ile Trp Leu Gly Lys Tyr Gly Asp Ile Gly
 130 135 140
 Pro Ile Gly Ser Ser Gln Gly Thr Val Asn Val Gly Gly Gln Ser Trp
 145 150 155 160
 Thr Leu Tyr Tyr Gly Tyr Asn Gly Ala Met Gln Val Tyr Ser Phe Val
 165 170 175
 Ala Gln Thr Asn Thr Thr Asn Tyr Ser Gly Asp Val Lys Asn Phe Phe
 180 185 190
 Asn Tyr Leu Arg Asp Asn Lys Gly Tyr Asn Ala Ala Gly Gln Tyr Val
 195 200 205
 Leu Ser Tyr Gln Phe Gly Thr Glu Pro Phe Thr Gly Ser Gly Thr Leu

210
Asn Val Ala Ser Trp Thr Ala Ser Ile Asn
225 230

220

<210> 4
<211> 826
<212> DNA
<213> Trichoderma reesei

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tcagccgggt ctggatttgg ctgctgacg gcggtatcgc tcagcggcgg ggctcctgg 180
cacgcagact ggcagtgggc cggcggccag aacaacgtca agtcgtacca gaactctcag 240
attgccattc cccagaagag gaccgtcaac agcatcagca gcatgccac cactgccagc 300
tggagctaca gcgggagcaa catccgcgct aatgttgcgt atgacttggt caccgcagcc 360
aaccggaatc atgtcacgta ctcgggagac tacgaactca tgatctggta agccataaga 420
agtggacctc cttgatatgt tgcactaaca acatgtcttg aggcttggca aatacggcga 480
tattggggccg attgggtcct cacaggaac agtcaacgtc ggtggccaga gctggacgct 540
ctactatggc tacaacggag ccatgcaagt ctattccttt gtggccca ccaacactac 600
caactacagc ggagatgtca agaacttctt caattatctc cgagacaata aaggatacaa 660
cgctgcaggg caatatgttc ttagtaagtc accctcactg tgactgggct gagtttggtg 720
caacgtttgc taacaaaacc ttcgtatagg ctaccaattt ggtaccgagc ccttcacggg 780
cagtggaaact ctgaacgtcg catcctggac cgcattctac aactaa 826

<210> 5
<211> 222
<212> PRT
<213> Trichoderma reesei

<400> 5
Met Val Ser Phe Thr Ser Leu Leu Ala Ala Ser Pro Pro Ser Arg Ala
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Ser Cys Arg Pro Ala Ala Glu Val Glu Ser Val Ala Val Glu Lys Arg
20 25 30
Gln Thr Ile Gln Pro Gly Thr Gly Tyr Asn Asn Gly Tyr Phe Tyr Ser
35 40 45
Tyr Trp Asn Asp Gly His Gly Gly Val Thr Tyr Thr Asn Gly Pro Gly
50 55 60
Gly Gln Phe Ser Val Asn Trp Ser Asn Ser Gly Asn Phe Val Gly Gly
65 70 75 80
Lys Gly Trp Gln Pro Gly Thr Lys Asn Lys Val Ile Asn Phe Ser Gly
85 90 95
Ser Tyr Asn Pro Asn Gly Asn Ser Tyr Leu Ser Val Tyr Gly Trp Ser
100 105 110
Arg Asn Pro Leu Ile Glu Tyr Tyr Ile Val Glu Asn Phe Gly Thr Tyr
115 120 125
Asn Pro Ser Thr Gly Ala Thr Lys Leu Gly Glu Val Thr Ser Asp Gly
130 135 140
Ser Val Tyr Asp Ile Tyr Arg Thr Gln Arg Val Asn Gln Pro Ser Ile
145 150 155 160
Ile Gly Thr Ala Thr Phe Tyr Gln Tyr Trp Ser Val Arg Arg Asn His
165 170 175
Arg Ser Ser Gly Ser Val Asn Thr Ala Asn His Phe Asn Ala Trp Ala
180 185 190
Gln Gln Gly Leu Thr Leu Gly Thr Met Asp Tyr Gln Ile Val Ala Val
195 200 205

Glu Gly Tyr Phe Ser Ser Gly Ser Ala Ser Ile Thr Val Ser
 210 215 220

<210> 6
 <211> 227
 <212> PRT
 <213> Humicola insolens

<400> 6
 Met Val Ser Leu Lys Ser Val Leu Ala Ala Ala Thr Ala Val Ser Ser
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 Ala Ile Ala Ala Pro Phe Asp Phe Val Pro Arg Asp Asn Ser Thr Ala
 20 25 30
 Leu Gln Ala Arg Gln Val Thr Pro Asn Ala Glu Gly Trp His Asn Gly
 35 40 45
 Tyr Phe Tyr Ser Trp Trp Ser Asp Gly Gly Gly Gln Val Gln Tyr Thr
 50 55 60
 Asn Leu Glu Gly Ser Arg Tyr Gln Val Arg Trp Arg Asn Thr Gly Asn
 65 70 75 80
 Phe Val Gly Gly Lys Gly Trp Asn Pro Gly Thr Gly Arg Thr Ile Asn
 85 90 95
 Tyr Gly Gly Tyr Phe Asn Pro Gln Gly Asn Gly Tyr Leu Ala Val Tyr
 100 105 110
 Gly Trp Thr Arg Asn Pro Leu Val Glu Tyr Tyr Val Ile Glu Ser Tyr
 115 120 125
 Gly Thr Tyr Asn Pro Gly Ser Gln Ala Gln Tyr Lys Gly Thr Phe Tyr
 130 135 140
 Thr Asp Gly Asp Gln Tyr Asp Ile Phe Val Ser Thr Arg Tyr Asn Gln
 145 150 155 160
 Pro Ser Ile Asp Gly Thr Arg Thr Phe Gln Tyr Trp Ser Ile Arg
 165 170 175
 Lys Asn Lys Arg Val Gly Gly Ser Val Asn Met Gln Asn His Phe Asn
 180 185 190
 Ala Trp Gln Gln His Gly Met Pro Leu Gly Gln His Tyr Tyr Gln Val
 195 200 205
 Val Ala Thr Glu Gly Tyr Gln Ser Ser Gly Glu Ser Asp Ile Tyr Val
 210 215 220
 Gln Thr His
 225

<210> 7
 <211> 210
 <212> PRT
 <213> Bacillus stearothermophilus

<400> 7
 Met Lys Leu Lys Lys Lys Met Leu Thr Leu Leu Leu Thr Ala Ser Met
 1 5 10 15
 Ser Phe Gly Leu Phe Gly Ala Thr Ser Ser Ala Ala Thr Asp Tyr Trp
 20 25 30
 Gln Tyr Trp Thr Asp Gly Gly Gly Met Val Asn Ala Val Asn Gly Pro
 35 40 45
 Gly Gly Asn Tyr Ser Val Thr Trp Gln Asn Thr Gly Asn Phe Val Val
 50 55 60
 Gly Lys Gly Trp Thr Val Gly Ser Pro Asn Arg Val Ile Asn Tyr Asn
 65 70 75 80
 Ala Gly Ile Trp Glu Pro Ser Gly Asn Gly Tyr Leu Thr Leu Tyr Gly

				85					90					95			
Trp	Thr	Arg	Asn	Ala	Leu	Ile	Glu	Tyr	Tyr	Val	Val	Asp	Ser	Trp	Gly		
			100					105					110				
Thr	Tyr	Arg	Pro	Thr	Gly	Asn	Tyr	Lys	Gly	Thr	Val	Asn	Ser	Asp	Gly		
		115					120					125					
Gly	Thr	Tyr	Asp	Ile	Tyr	Thr	Thr	Met	Arg	Tyr	Asn	Ala	Pro	Ser	Ile		
	130					135					140						
Asp	Gly	Thr	Gln	Thr	Phe	Gln	Gln	Phe	Trp	Ser	Val	Arg	Gln	Ser	Lys		
145					150					155					160		
Arg	Pro	Thr	Gly	Ser	Asn	Val	Ser	Ile	Thr	Phe	Ser	Asn	His	Val	Asn		
			165					170					175				
Ala	Trp	Arg	Ser	Lys	Gly	Met	Asn	Leu	Gly	Ser	Ser	Trp	Ala	Tyr	Gln		
		180						185					190				
Val	Leu	Ala	Thr	Glu	Gly	Tyr	Gln	Ser	Ser	Gly	Arg	Ser	Asn	Val	Thr		
	195						200					205					
Val	Trp																
	210																

<210> 8

<211> 229

<212> PRT

<213> Trichoderma reesei

<400> 8

Met	Val	Ala	Phe	Ser	Ser	Leu	Ile	Cys	Ala	Leu	Thr	Ser	Ile	Ala	Ser		
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Thr	Leu	Ala	Met	Pro	Thr	Gly	Leu	Glu	Pro	Glu	Ser	Ser	Val	Asn	Val		
		20						25					30				
Thr	Glu	Arg	Gly	Met	Tyr	Asp	Phe	Val	Leu	Gly	Ala	His	Asn	Asp	His		
	35					40						45					
Arg	Arg	Arg	Ala	Ser	Ile	Asn	Tyr	Asp	Gln	Asn	Tyr	Gln	Thr	Gly	Gly		
	50				55					60							
Gln	Val	Ser	Tyr	Ser	Pro	Ser	Asn	Thr	Gly	Phe	Ser	Val	Asn	Trp	Asn		
65				70					75						80		
Thr	Gln	Asp	Asp	Phe	Val	Val	Gly	Val	Gly	Trp	Thr	Thr	Gly	Ser	Ser		
			85					90					95				
Ala	Pro	Ile	Asn	Phe	Gly	Gly	Ser	Phe	Ser	Val	Asn	Ser	Gly	Thr	Gly		
		100						105					110				
Leu	Leu	Ser	Val	Tyr	Gly	Trp	Ser	Thr	Asn	Pro	Leu	Val	Glu	Tyr	Tyr		
	115					120						125					
Ile	Met	Glu	Asp	Asn	His	Asn	Tyr	Pro	Ala	Gln	Gly	Thr	Val	Lys	Gly		
	130			135							140						
Thr	Val	Thr	Ser	Asp	Gly	Ala	Thr	Tyr	Thr	Ile	Trp	Glu	Asn	Thr	Arg		
145				150				155						160			
Val	Asn	Glu	Pro	Ser	Ile	Gln	Gly	Thr	Ala	Thr	Phe	Asn	Gln	Tyr	Ile		
			165					170					175				
Ser	Val	Arg	Asn	Ser	Pro	Arg	Thr	Ser	Gly	Thr	Val	Thr	Val	Gln	Asn		
		180						185					190				
His	Phe	Asn	Ala	Trp	Ala	Ser	Leu	Gly	Leu	His	Leu	Gly	Gln	Met	Asn		
	195					200						205					
Tyr	Gln	Val	Val	Ala	Val	Glu	Gly	Trp	Gly	Gly	Ser	Gly	Ser	Ala	Ser		
	210				215						220						
Gln	Ser	Val	Ser	Asn													
	225																

<210> 9

<211> 211

<212> PRT

<213> *Aspergillus awamori*

<400> 9

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Met Lys Val Thr Ala Ala Phe Ala Gly Leu Leu Val Thr Ala Phe Ala
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Ala Pro Val Pro Glu Pro Val Leu Val Ser Arg Ser Ala Gly Ile Asn
          20          25          30
Tyr Val Gln Asn Tyr Asn Gly Asn Leu Gly Asp Phe Thr Tyr Asp Glu
          35          40          45
Ser Ala Gly Thr Phe Ser Met Tyr Trp Glu Asp Gly Val Ser Ser Asp
          50          55          60
Phe Val Val Gly Leu Gly Trp Thr Thr Gly Ser Ser Asn Ala Ile Thr
65          70          75          80
Tyr Ser Ala Glu Tyr Ser Ala Ser Gly Ser Ser Ser Tyr Leu Ala Val
          85          90          95
Tyr Gly Trp Val Asn Tyr Pro Gln Ala Glu Tyr Tyr Ile Val Glu Asp
          100          105          110
Tyr Gly Asp Tyr Asn Pro Cys Ser Ser Ala Thr Ser Leu Gly Thr Val
          115          120          125
Tyr Ser Asp Gly Ser Thr Tyr Gln Val Cys Thr Asp Thr Arg Thr Asn
          130          135          140
Glu Pro Ser Ile Thr Gly Thr Ser Thr Phe Thr Gln Tyr Phe Ser Val
145          150          155          160
Arg Glu Ser Thr Arg Thr Ser Gly Thr Val Thr Val Ala Asn His Phe
          165          170          175
Asn Phe Trp Ala Gln His Gly Phe Gly Asn Ser Asp Phe Asn Tyr Gln
          180          185          190
Val Met Ala Val Glu Ala Trp Ser Gly Ala Gly Ser Ala Ser Val Thr
          195          200          205
Ile Ser Ser
          210
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<210> 10

<211> 330

<212> PRT

<213> *Bacillus stearothermophilus*

<400> 10

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Arg Ile Gly Ala Ala Val Asn Pro Val Thr Leu Glu Ala Gln Gln Ser
          20          25          30
Leu Leu Ile Arg His Val Asn Ser Leu Thr Ala Glu Asn His Met Lys
          35          40          45
Phe Glu His Leu Gln Pro Glu Glu Gly Arg Phe Thr Phe Asp Ile Ala
          50          55          60
Ile Lys Ser Ser Thr Ser Pro Phe Ser Ser His Gly Val Arg Gly His
65          70          75          80
Thr Leu Val Trp His Asn Gln Thr Pro Ser Trp Val Phe Gln Asp Ser
          85          90          95
Gln Gly His Phe Val Gly Arg Asp Val Leu Leu Glu Arg Met Lys Ser
          100          105          110
His Ile Ser Thr Val Val Gln Arg Tyr Lys Gly Lys Val Tyr Cys Trp
          115          120          125
Asp Val Ile Asn Glu Ala Val Ala Asp Glu Gly Ser Glu Trp Leu Arg
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130		135		140
Ser Ser Thr Trp Arg Gln Ile Ile Gly Asp Asp Phe Ile Gln Gln Ala				
145		150		155
Phe Leu Tyr Ala His Glu Ala Asp Pro Glu Ala Leu Leu Phe Tyr Asn				160
	165		170	175
Asp Tyr Asn Glu Cys Phe Pro Glu Lys Arg Glu Lys Ile Tyr Thr Leu				
	180		185	190
Val Lys Ser Leu Arg Asp Lys Gly Ile Pro Ile His Gly Ile Gly Met				
	195		200	205
Gln Ala His Trp Ser Leu Asn Arg Pro Thr Leu Asp Glu Ile Arg Ala				
	210		215	220
Ala Ile Glu Arg Tyr Ala Ser Leu Gly Val Ile Leu His Ile Thr Glu				
225		230		235
Leu Asp Ile Ser Met Phe Glu Phe Asp Asp His Arg Lys Asp Leu Ala				240
	245		250	255
Ala Pro Thr Asn Glu Met Val Glu Arg Gln Ala Glu Arg Tyr Glu Gln				
	260		265	270
Ile Phe Ser Leu Phe Lys Glu Tyr Arg Asp Val Ile Gln Asn Val Thr				
	275		280	285
Phe Trp Gly Ile Ala Asp Asp His Thr Trp Leu Asp His Phe Pro Val				
	290		295	300
Gln Gly Arg Lys Asn Trp Pro Leu Leu Phe Asp Glu Gln His Asn Pro				
305		310		315
Lys Pro Ala Phe Trp Arg Val Val Asn Ile				320
	325		330	

<210> 11

<211> 190

<212> PRT

<213> *Trichoderma reesei*

<400> 11

Gln Thr Ile Gln Pro Gly Thr Gly Tyr Asn Asn Gly Tyr Phe Tyr Ser	
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Tyr Trp Asn Asp Gly His Gly Gly Val Thr Tyr Thr Asn Gly Pro Gly	
	20
Gly Gln Phe Ser Val Asn Trp Ser Asn Ser Gly Asn Phe Val Gly Gly	
	35
Lys Gly Trp Gln Pro Gly Thr Lys Asn Lys Val Ile Asn Phe Ser Gly	
	50
Ser Tyr Asn Pro Asn Gly Asn Ser Tyr Leu Ser Val Tyr Gly Trp Ser	
65	70
Arg Asn Pro Leu Ile Glu Tyr Tyr Ile Val Glu Asn Phe Gly Thr Tyr	
	85
Asn Pro Ser Thr Gly Ala Thr Lys Leu Gly Glu Val Thr Ser Asp Gly	
	100
Ser Val Tyr Asp Ile Tyr Arg Thr Gln Arg Val Asn Gln Pro Ser Ile	
	115
Ile Gly Thr Ala Thr Phe Tyr Gln Tyr Trp Ser Val Arg Arg Asn His	
	130
Arg Ser Ser Gly Ser Val Asn Thr Ala Asn His Phe Asn Ala Trp Ala	
145	150
Gln Gln Gly Leu Thr Leu Gly Thr Met Asp Tyr Gln Ile Val Ala Val	
	165
Glu Gly Tyr Phe Ser Ser Gly Ser Ala Ser Ile Thr Val Ser	
	180

<210> 12
 <211> 237
 <212> PRT
 <213> *Aspergillus awamori*

<400> 12
 Met Lys Ala Phe His Leu Leu Ala Ala Leu Ser Gly Ala Ala Val Ala
 1 5 10 15
 Gln Gln Ala Gln Leu Cys Asp Gln Tyr Ala Thr Tyr Thr Gly Gly Val
 20 25 30
 Tyr Thr Ile Asn Asn Asn Leu Trp Gly Lys Asp Ala Gly Ser Gly Ser
 35 40 45
 Gln Cys Thr Thr Val Asn Ser Ala Ser Ser Ala Gly Thr Ser Trp Ser
 50 55 60
 Thr Lys Trp Asn Trp Ser Gly Gly Glu Asn Ser Val Lys Ser Tyr Ala
 65 70 75 80
 Asn Ser Gly Leu Ser Phe Asn Lys Lys Leu Val Ser Gln Ile Ser His
 85 90 95
 Ile Pro Thr Ala Ala Arg Trp Ser Tyr Asp Asn Thr Cys Ile Arg Arg
 100 105 110
 Gly Arg Ala Tyr Asp Leu Phe Thr Ala Ala Asp Ile Asn His Val Thr
 115 120 125
 Trp Ser Gly Asp Tyr Glu Leu Met Ile Trp Leu Ala Arg Tyr Gly Gly
 130 135 140
 Val Gln Pro Leu Gly Ser Gln Ile Ala Thr Ala Thr Val Glu Gly Gln
 145 150 155 160
 Thr Trp Glu Leu Trp Tyr Gly Val Asn Gly Ala Gln Lys Thr Tyr Ser
 165 170 175
 Phe Val Ala Ala Asn Pro Ile Thr Ser Phe Gln Gly Asp Ile Asn Asp
 180 185 190
 Phe Phe Lys Tyr Leu Thr Gln Asn His Gly Phe Pro Ala Ser Ser Gln
 195 200 205
 Tyr Leu Ile Thr Leu Gln Phe Gly Thr Glu Pro Phe Thr Gly Gly Pro
 210 215 220
 Ala Thr Leu Asn Val Ala Asp Trp Ser Ala Ser Val Gln
 225 230 235

<210> 13
 <211> 233
 <212> PRT
 <213> *Trichoderma viride*

<400> 13
 Met Lys Phe Leu Gln Ile Ala Pro Thr Leu Leu Pro Val Ala Leu Ala
 1 5 10 15
 Gln Ser Ser Cys Ser Gln Tyr Ala Thr Phe Ser Gly Gly Asn Tyr Ala
 20 25 30
 Leu Ser Asn Asn Leu Trp Gly Gln Ser Ala Gly Ser Gly Ser Gly Cys
 35 40 45
 Ile Thr Asp Val Ser Leu Gly Gly Ser Ala Val Trp Ser Thr Thr Trp
 50 55 60
 Asp Trp Ser Gly Gly Gln Ser Asn Val Lys Gly Tyr Pro Asn Ile Ala
 65 70 75 80
 Leu Asn Ile Pro Asn Lys Arg Leu Val Ser Ser Ile Ser Ser Met Pro
 85 90 95
 Thr Thr Ala Gln Trp Ser Tyr Ser Gly Ser Ser Ile Arg Ala Asp Val

			100						105				110			
Ala	Tyr	Asp	Leu	Phe	Thr	Ala	Ser	Asn	Pro	Asn	His	Val	Thr	Tyr	Ser	
		115						120					125			
Gly	Asp	Tyr	Glu	Leu	Met	Ile	Trp	Leu	Gly	Lys	Tyr	Gly	Asp	Ile	Gln	
	130					135					140					
Pro	Ile	Gly	Ser	Ser	Gln	Gly	Thr	Val	Asn	Val	Gly	Gly	Thr	Ser	Trp	
145					150					155					160	
Asn	Leu	Trp	Tyr	Gly	Pro	Asn	Gly	Ser	Met	Gln	Val	Tyr	Ser	Phe	Val	
				165					170					175		
Ala	Pro	Gly	Asn	Leu	Thr	Asn	Trp	Ser	Gly	Asp	Val	Lys	Asn	Phe	Tyr	
		180						185					190			
Thr	Tyr	Leu	Gln	Asn	Asn	Lys	Gly	Tyr	Pro	Ala	Ser	Ser	Gln	Tyr	Val	
	195					200						205				
Leu	Ser	Tyr	Gln	Phe	Gly	Thr	Glu	Ala	Phe	Thr	Gly	Ser	Gly	Thr	Leu	
	210					215					220					
Asn	Asn	Thr	Trp	Thr	Ala	Ser	Ile	Asn								
225					230											

<210> 14
 <211> 234
 <212> PRT
 <213> Hypocrea koningii

Met	Lys	Leu	Ile	His	Val	Leu	Pro	Ala	Leu	Ile	Pro	Ala	Ala	Leu	Ala	
1				5					10					15		
Gln	Thr	Ser	Cys	Asp	Gln	Tyr	Ala	Val	Phe	Thr	Gly	Ser	Asp	Tyr	Thr	
			20					25					30			
Val	Ser	Asn	Asn	Leu	Trp	Gly	Gln	Ser	Ala	Gly	Ser	Gly	Phe	Gly	Cys	
	35					40						45				
Val	Thr	Ala	Glu	Ser	Leu	Ser	Gly	Ser	Ala	Ser	Trp	His	Ala	Asp	Trp	
	50					55					60					
Gln	Trp	Ser	Gly	Gly	Gln	Asn	Asn	Val	Lys	Ser	Tyr	Gln	Asn	Ser	Gln	
65					70					75					80	
Ile	Pro	Ile	Pro	Gln	Lys	Arg	Thr	Val	Asn	Ser	Ile	Ser	Ser	Met	Pro	
				85					90					95		
Thr	Thr	Ala	Ser	Trp	Ser	Tyr	Thr	Gly	Ser	Asp	Ile	Arg	Ala	Asn	Val	
			100					105					110			
Ala	Tyr	Asp	Leu	Phe	Thr	Ala	Ala	Asn	Pro	Asn	His	Val	Thr	Tyr	Ser	
	115					120						125				
Gly	Asp	Tyr	Glu	Leu	Met	Ile	Trp	Leu	Gly	Arg	Tyr	Gly	Asp	Ile	Gly	
	130					135					140					
Pro	Ile	Gly	Ser	Ser	Gln	Gly	Thr	Val	Asn	Val	Gly	Gly	Gln	Ser	Trp	
145					150					155					160	
Thr	Leu	Tyr	Tyr	Gly	Tyr	Asn	Gly	Ala	Met	Gln	Val	Tyr	Ser	Phe	Val	
				165				170						175		
Ala	Gln	Thr	Asn	Thr	Thr	Ser	Tyr	Ser	Gly	Asp	Val	Lys	Asn	Phe	Phe	
		180						185					190			
Asn	Tyr	Leu	Arg	Asp	Asn	Lys	Gly	Tyr	Asn	Ala	Ala	Gly	Gln	Tyr	Val	
	195					200						205				
Leu	Ser	Tyr	Gln	Phe	Gly	Thr	Glu	Pro	Phe	Thr	Gly	Ser	Gly	Thr	Leu	
	210					215					220					
Asn	Val	Ala	Ser	Trp	Thr	Ala	Ser	Ile	Asn							
225					230											

<210> 15
 <211> 234

<212> PRT

<213> *Hypocrea schweinitzii*

<400> 15

Met	Lys	Phe	Leu	Gln	Val	Leu	Pro	Ala	Ile	Leu	Pro	Ala	Ala	Leu	Ala
1				5					10					15	
Gln	Thr	Ser	Cys	Asp	Gln	Tyr	Ala	Thr	Phe	Ser	Gly	Asn	Gly	Tyr	Ile
			20					25					30		
Val	Ser	Asn	Asn	Leu	Trp	Gly	Ala	Ser	Ala	Gly	Ser	Gly	Phe	Gly	Cys
		35					40					45			
Val	Thr	Ser	Val	Ser	Leu	Asn	Gly	Ala	Ala	Ser	Trp	His	Ala	Asp	Trp
	50					55					60				
Gln	Trp	Ser	Gly	Gly	Gln	Asn	Asn	Val	Lys	Ser	Tyr	Gln	Asn	Val	Gln
65					70					75					80
Ile	Asn	Ile	Pro	Gln	Lys	Arg	Thr	Val	Asn	Ser	Ile	Gly	Ser	Met	Pro
				85					90					95	
Thr	Thr	Ala	Ser	Trp	Ser	Tyr	Ser	Gly	Ser	Asp	Ile	Arg	Ala	Asn	Val
			100					105					110		
Ala	Tyr	Asp	Leu	Phe	Thr	Ala	Ala	Asn	Pro	Asn	His	Val	Thr	Tyr	Ser
		115					120					125			
Gly	Asp	Tyr	Glu	Leu	Met	Ile	Trp	Leu	Gly	Lys	Tyr	Gly	Asp	Ile	Gly
	130					135					140				
Pro	Ile	Gly	Ser	Ser	Gln	Gly	Thr	Val	Asn	Val	Gly	Gly	Gln	Thr	Trp
145					150					155					160
Thr	Leu	Tyr	Tyr	Gly	Tyr	Asn	Gly	Ala	Met	Gln	Val	Tyr	Ser	Phe	Val
				165				170						175	
Ala	Gln	Ser	Asn	Thr	Thr	Ser	Tyr	Ser	Gly	Asp	Val	Lys	Asn	Phe	Phe
			180					185					190		
Asn	Tyr	Leu	Arg	Asp	Asn	Lys	Gly	Tyr	Asn	Ala	Gly	Gly	Gln	Tyr	Val
	195						200					205			
Leu	Ser	Tyr	Gln	Phe	Gly	Thr	Glu	Pro	Phe	Thr	Gly	Ser	Gly	Thr	Leu
	210					215					220				
Asn	Val	Ala	Ser	Trp	Thr	Ala	Ser	Ile	Asn						
225					230										

<210> 16

<211> 237

<212> PRT

<213> *Stachybotrys echinata*

<400> 16

Met	Lys	Val	Ala	Ala	Leu	Leu	Val	Ala	Leu	Ser	Pro	Leu	Ala	Phe	Ala
1				5					10					15	
Gln	Ser	Leu	Cys	Asp	Gln	Tyr	Ser	Tyr	Tyr	Ser	Ser	Asn	Gly	Tyr	Glu
			20					25					30		
Phe	Asn	Asn	Asn	Met	Trp	Gly	Arg	Asn	Ser	Gly	Gln	Gly	Asn	Gln	Cys
		35					40					45			
Thr	Tyr	Val	Asp	Tyr	Ser	Ser	Pro	Asn	Gly	Val	Gly	Trp	Arg	Val	Asn
	50					55					60				
Trp	Asn	Trp	Ser	Gly	Gly	Asp	Asn	Asn	Val	Lys	Ser	Tyr	Pro	Tyr	Ser
65					70					75					80
Gly	Arg	Gln	Leu	Pro	Thr	Lys	Arg	Ile	Val	Ser	Trp	Ile	Gly	Ser	Leu
				85					90					95	
Pro	Thr	Thr	Val	Ser	Trp	Asn	Tyr	Gln	Gly	Asn	Asn	Leu	Arg	Ala	Asn
			100					105					110		
Val	Ala	Tyr	Asp	Leu	Phe	Thr	Ala	Ala	Asn	Pro	Asn	His	Pro	Asn	Ser
		115					120					125			

Ser Gly Asp Tyr Glu Leu Met Ile Trp Leu Gly Arg Leu Gly Asn Val
 130 135 140
 Tyr Pro Ile Gly Asn Gln Val Ala Thr Val Asn Ile Ala Gly Gln Gln
 145 150 155 160
 Trp Asn Leu Tyr Tyr Gly Tyr Asn Gly Ala Met Gln Val Tyr Ser Phe
 165 170 175
 Val Ser Pro Asn Gln Leu Asn Tyr Phe Ser Gly Asn Val Lys Asp Phe
 180 185 190
 Phe Thr Tyr Leu Gln Tyr Asn Arg Ala Tyr Pro Ala Asp Ser Gln Tyr
 195 200 205
 Leu Ile Thr Tyr Gln Phe Gly Thr Glu Pro Phe Thr Gly Gln Asn Ala
 210 215 220
 Val Phe Thr Val Ser Asn Trp Ser Ala Gln Gln Asn Asn
 225 230 235

<210> 17

<211> 238

<212> PRT

<213> *Fusarium equiseti*

<400> 17

Met Lys Ser Thr Leu Leu Leu Ala Gly Ala Phe Ala Pro Leu Ala Phe
 1 5 10 15
 Ala Lys Asp Leu Cys Glu Gln Tyr Gly Tyr Leu Ser Ser Asp Gly Tyr
 20 25 30
 Ser Leu Asn Asn Asn Val Trp Gly Lys Asp Ser Gly Thr Gly Asp Gln
 35 40 45
 Cys Thr His Val Asn Trp Asn Asn Ala Asn Gly Ala Gly Trp Asp Val
 50 55 60
 Glu Trp Asn Trp Ser Gly Gly Lys Asp Asn Val Lys Ser Tyr Pro Asn
 65 70 75 80

Ser Ala Leu Leu Ile Gly Glu Asp Lys Lys Thr Ile Ser Ser Ile Thr
 85 90 95
 Asn Met Gln Ser Thr Ala Glu Trp Lys Tyr Ser Gly Asp Asn Leu Arg
 100 105 110
 Ala Asp Val Ala Tyr Asp Leu Phe Thr Ala Ala Asp Pro Asn His Glu
 115 120 125
 Thr Ser Ser Gly Glu Tyr Glu Leu Met Val Trp Leu Ala Arg Ile Gly
 130 135 140
 Gly Val Gln Pro Ile Gly Ser Leu Gln Thr Ser Val Thr Ile Glu Gly
 145 150 155 160
 His Thr Trp Glu Leu Trp Val Gly Met Asn Gly Ser Met Lys Val Phe
 165 170 175
 Ser Phe Val Ala Pro Thr Pro Val Asn Asn Phe Asn Ala Asp Ile Lys
 180 185 190
 Gln Phe Trp Asp Tyr Leu Thr Lys Ser Gln Asn Phe Pro Ala Asp Asn
 195 200 205
 Gln Tyr Leu Leu Thr Phe Gln Phe Gly Thr Glu Pro Phe Thr Gly Asp
 210 215 220
 Asn Ala Lys Phe Thr Val Thr Asn Phe Asn Ala His Leu Lys
 225 230 235

<210> 18

<211> 237

<212> PRT

<213> *Bionectria ochroleuca*

<400> 18

Met	Lys	Thr	Gly	Ile	Ala	Tyr	Leu	Ala	Ala	Val	Leu	Pro	Leu	Ala	Met
1				5					10					15	
Ala	Glu	Ser	Leu	Cys	Asp	Gln	Tyr	Ala	Tyr	Leu	Ser	Arg	Asp	Gly	Tyr
			20					25					30		
Asn	Phe	Asn	Asn	Asn	Glu	Trp	Gly	Ala	Ala	Thr	Gly	Thr	Gly	Asp	Gln
		35					40					45			
Cys	Thr	Tyr	Val	Asp	Ser	Thr	Ser	Ser	Gly	Gly	Val	Ser	Trp	His	Ser
	50					55					60				
Asp	Trp	Thr	Asn	Ser	Gly	Ser	Glu	Ser	Glu	Ile	Lys	Ser	Tyr	Pro	Tyr
65					70					75					80
Ser	Gly	Leu	Asp	Leu	Pro	Glu	Lys	Lys	Ile	Val	Thr	Ser	Ile	Gly	Ser
				85					90					95	
Ile	Ser	Thr	Gly	Ala	Glu	Trp	Ser	Tyr	Ser	Gly	Ser	Asn	Ile	Arg	Ala
			100					105					110		
Asp	Val	Ala	Tyr	Asp	Ile	Phe	Thr	Ala	Ala	Asp	Pro	Asn	His	Ala	Thr
		115					120					125			
Ser	Ser	Gly	Asp	Tyr	Glu	Val	Met	Ile	Trp	Leu	Ala	Asn	Leu	Gly	Gly
		130				135					140				
Leu	Thr	Pro	Ile	Gly	Ser	Pro	Ile	Gly	Thr	Val	Lys	Ala	Ala	Gly	Arg
145				150						155					160
Asp	Trp	Glu	Leu	Trp	Asp	Gly	Tyr	Asn	Gly	Ala	Met	Arg	Val	Tyr	Ser
				165					170					175	
Phe	Val	Ala	Pro	Ser	Gln	Leu	Asn	Ser	Phe	Asp	Gly	Glu	Ile	Met	Asp
		180					185					190			
Phe	Phe	Tyr	Val	Val	Lys	Asp	Met	Arg	Gly	Phe	Pro	Ala	Asp	Ser	Gln
		195				200					205				
His	Leu	Leu	Thr	Val	Gln	Phe	Gly	Thr	Glu	Pro	Ile	Ser	Gly	Ser	Gly
	210				215					220					
Ala	Lys	Phe	Ser	Val	Ser	His	Trp	Ser	Ala	Lys	Leu	Gly			
225					230					235					

<210> 19

<211> 236

<212> PRT

<213> Bionectria ochroleuca

<400> 19

Met	Lys	Phe	Gln	Leu	Leu	Ser	Leu	Thr	Ala	Phe	Ala	Pro	Leu	Ser	Leu
1				5					10					15	
Ala	Ala	Leu	Cys	Gly	Gln	Tyr	Gln	Ser	Gln	Ser	Gln	Gly	Gly	Tyr	Ile
			20					25					30		
Phe	Asn	Asn	Asn	Lys	Trp	Gly	Gln	Gly	Ser	Gly	Ser	Gly	Ser	Gln	Cys
		35					40					45			
Leu	Thr	Ile	Asp	Lys	Thr	Trp	Asp	Ser	Asn	Val	Ala	Phe	His	Ala	Asp
	50					55					60				
Trp	Ser	Trp	Ser	Gly	Gly	Thr	Asn	Asn	Val	Lys	Ser	Tyr	Pro	Asn	Ala
65					70					75					80
Gly	Leu	Glu	Phe	Ser	Arg	Gly	Lys	Lys	Val	Ser	Ser	Ile	Gly	Thr	Ile
				85					90					95	
Asn	Gly	Gly	Ala	Asp	Trp	Asp	Tyr	Ser	Gly	Ser	Asn	Ile	Arg	Ala	Asn
			100					105					110		
Val	Ala	Tyr	Asp	Ile	Phe	Thr	Ser	Ala	Asp	Pro	Asn	His	Val	Thr	Ser
		115					120					125			
Ser	Gly	Asp	Tyr	Glu	Leu	Met	Ile	Trp	Leu	Gly	Lys	Leu	Gly	Asp	Ile
	130					135					140				

Tyr	Pro	Ile	Gly	Asn	Ser	Ile	Gly	Arg	Val	Lys	Ala	Ala	Asn	Arg	Glu
145					150					155					160
Trp	Asp	Leu	His	Val	Gly	Tyr	Asn	Gly	Ala	Met	Lys	Val	Phe	Ser	Phe
				165					170					175	
Val	Ala	Pro	Ser	Pro	Val	Thr	Arg	Phe	Asp	Gly	Asn	Ile	Met	Asp	Phe
			180					185					190		
Phe	Tyr	Val	Met	Arg	Asp	Met	Gln	Gly	Tyr	Pro	Met	Asp	Lys	Gln	Tyr
		195					200					205			
Leu	Leu	Thr	Leu	Gln	Phe	Gly	Thr	Glu	Pro	Phe	Thr	Gly	Ser	Asn	Ala
		210				215					220				
Lys	Phe	Ser	Cys	Trp	Tyr	Phe	Gly	Ala	Lys	Ile	Lys				
225					230					235					

<210> 20

<211> 240

<212> PRT

<213> Bionectria ochroleuca

<400> 20

Met	Lys	Ala	Asn	Ile	Val	Ile	Leu	Ser	Leu	Phe	Ala	Pro	Leu	Ala	Ala
1				5					10					15	
Val	Ala	Gln	Thr	Leu	Cys	Gly	Gln	Tyr	Ser	Ser	Asn	Thr	Gln	Gly	Gly
			20					25					30		
Tyr	Ile	Phe	Asn	Asn	Asn	Met	Trp	Gly	Met	Gly	Ser	Gly	Ser	Gly	Ser
		35					40					45			
Gln	Cys	Thr	Tyr	Val	Asp	Lys	Val	Trp	Ala	Glu	Gly	Val	Ala	Trp	His
	50					55					60				
Thr	Asp	Trp	Ser	Trp	Ser	Gly	Gly	Asp	Asn	Asn	Val	Lys	Ser	Tyr	Pro
65					70				75					80	
Tyr	Ser	Gly	Arg	Glu	Leu	Gly	Thr	Lys	Arg	Ile	Val	Ser	Ser	Ile	Lys
				85				90						95	
Ser	Ile	Ser	Ser	Gly	Ala	Asp	Trp	Asp	Tyr	Thr	Gly	Ser	Asn	Leu	Arg
			100					105					110		
Ala	Asn	Ala	Ala	Tyr	Asp	Ile	Phe	Thr	Ser	Ala	Asn	Pro	Asn	His	Ala
		115					120					125			
Thr	Ser	Ser	Gly	Asp	Tyr	Glu	Val	Met	Ile	Trp	Leu	Gly	Arg	Tyr	Gly
		130				135					140				
Gly	Val	Tyr	Pro	Ile	Gly	Asn	Ser	Ile	Gly	Thr	Val	Arg	Ala	Ala	Gly
145					150					155					160
Arg	Asp	Trp	Ala	Leu	His	Ile	Gly	Tyr	Asn	Gly	Ala	Met	Lys	Val	Phe
				165					170					175	
Ser	Phe	Val	Ala	Ala	Asn	Pro	Val	Thr	Arg	Phe	Asp	Gly	Glu	Ile	Met
			180					185					190		
Asp	Phe	Phe	Tyr	Leu	Leu	Arg	Asp	Met	Gln	Gly	Tyr	Pro	Met	Thr	Ser
		195					200					205			
Gln	Tyr	Leu	Leu	Thr	Leu	Gln	Phe	Gly	Thr	Glu	Pro	Phe	Thr	Gly	Ser
	210					215					220				
Gly	Ala	Lys	Phe	Asn	Cys	Trp	Tyr	Phe	Gly	Ala	Thr	Leu	Ser	Tyr	Trp
225					230					235					240

<210> 21

<211> 254

<212> PRT

<213> Humicola grisea

<400> 21

Met Leu Lys Ser Ala Leu Leu Leu Gly Ala Ala Ala Val Ser Val Gln

1				5					10					15			
Ser	Ala	Ser	Ile	Pro	Thr	Ile	Pro	Ala	Asn	Leu	Glu	Pro	Arg	Gln	Ile		
			20					25					30				
Arg	Ser	Leu	Cys	Glu	Leu	Tyr	Gly	Tyr	Trp	Ser	Gly	Asn	Gly	Tyr	Glu		
		35					40					45					
Leu	Leu	Asn	Asn	Leu	Trp	Gly	Lys	Asp	Thr	Ala	Thr	Ser	Gly	Trp	Gln		
		50				55					60						
Cys	Thr	Tyr	Leu	Asp	Gly	Thr	Asn	Asn	Gly	Gly	Ile	Gln	Trp	Asn	Thr		
65				70					75					80			
Ala	Trp	Glu	Trp	Gln	Gly	Ala	Pro	Asp	Asn	Val	Lys	Asn	Tyr	Pro	Tyr		
			85					90					95				
Val	Gly	Lys	Gln	Ile	Gln	Arg	Gly	Arg	Lys	Ile	Ser	Asp	Ile	Asn	Ser		
			100					105					110				
Met	Arg	Thr	Ser	Val	Ser	Trp	Thr	Tyr	Asp	Arg	Thr	Asp	Leu	Arg	Ala		
		115					120					125					
Asn	Val	Ala	Tyr	Asp	Val	Phe	Thr	Ala	Arg	Asp	Pro	Asp	His	Pro	Asn		
		130				135				140							
Trp	Gly	Gly	Asp	Tyr	Glu	Leu	Met	Ile	Trp	Leu	Ala	Arg	Tyr	Gly	Gly		
145				150					155					160			
Ile	Tyr	Pro	Ile	Gly	Thr	Phe	His	Ser	Gln	Val	Asn	Leu	Ala	Gly	Arg		
			165					170						175			
Thr	Trp	Asp	Leu	Trp	Thr	Gly	Tyr	Asn	Gly	Asn	Met	Arg	Val	Tyr	Ser		
		180					185					190					
Phe	Leu	Pro	Pro	Ser	Gly	Asp	Ile	Arg	Asp	Phe	Ser	Cys	Asp	Ile	Lys		
		195				200				205							
Asp	Phe	Phe	Asn	Tyr	Leu	Glu	Arg	Asn	His	Gly	Tyr	Pro	Ala	Arg	Glu		
		210				215				220							
Gln	Asn	Leu	Ile	Val	Tyr	Gln	Val	Gly	Thr	Glu	Cys	Phe	Thr	Gly	Gly		
225				230					235					240			
Pro	Ala	Arg	Phe	Thr	Cys	Arg	Asp	Phe	Arg	Ala	Asp	Leu	Trp				
			245					250									

<210> 22

<211> 247

<212> PRT

<213> Chaetomium brasiliense

<400> 22

Met	Lys	Leu	Thr	Leu	Val	Leu	Phe	Val	Ser	Ser	Leu	Ala	Ala	Ala	Thr		
1				5				10						15			
Pro	Leu	Gly	Trp	Arg	Glu	Arg	Arg	Gln	Gln	Val	Ser	Leu	Cys	Gly	Gln		
		20						25				30					
Ser	Ser	Ser	Trp	Ser	Gly	Asn	Gly	Tyr	Gln	Leu	Asn	Asn	Asn	Leu	Trp		
		35				40					45						
Gly	Gln	Ser	Arg	Ala	Thr	Ser	Gly	Ser	Gln	Cys	Thr	Tyr	Leu	Asp	Ser		
		50				55				60							
Ser	Ser	Asn	Ser	Gly	Ile	His	Trp	His	Thr	Thr	Trp	Thr	Trp	Glu	Gly		
65				70					75					80			
Gly	Glu	Gly	Glu	Val	Lys	Ser	Tyr	Ala	Tyr	Ser	Gly	Arg	Gln	Val	Ser		
			85					90					95				
Thr	Gly	Leu	Thr	Ile	Ala	Ser	Ile	Asp	Ser	Met	Gln	Thr	Ser	Val	Ser		
		100						105					110				
Trp	Glu	Tyr	Asn	Thr	Thr	Asp	Ile	Gln	Ala	Asn	Val	Ala	Tyr	Asp	Ile		
		115				120					125						
Phe	Thr	Ala	Glu	Asp	Pro	Asp	His	Glu	His	Ser	Ser	Gly	Asp	Tyr	Glu		
		130				135					140						
Val	Met	Ile	Trp	Leu	Ala	Arg	Tyr	Asn	Asn	Val	Ser	Pro	Ile	Gly	Ser		

145		150		155		160									
Ser	Val	Ala	Thr	Ala	Thr	Val	Gly	Gly	Asp	Thr	Trp	Asp	Leu	Phe	Ala
		165		170		175									
Gly	Ala	Asn	Gly	Asp	Met	Glu	Val	Tyr	Ser	Phe	Val	Ala	Glu	Asn	Thr
		180		185		190									
Met	Asn	Ser	Phe	Ser	Gly	Asp	Val	Lys	Asp	Phe	Phe	Asp	Tyr	Leu	Glu
		195		200		205									
Gln	Asn	Val	Gly	Phe	Pro	Val	Asp	Asp	Gln	Tyr	Leu	Leu	Val	Phe	Glu
		210		215		220									
Leu	Gly	Ser	Glu	Ala	Phe	Thr	Gly	Gly	Pro	Ala	Thr	Leu	Ser	Val	Ser
225				230		235									240
Gln	Phe	Ser	Ala	Asn	Ile	Ala									
				245											

<210> 23
 <211> 357
 <212> PRT
 <213> Bionectria ochroleuca

<400> 23
Met Lys Ser Ile Ile Ser Phe Phe Gly Leu Ala Thr Leu Val Ala Ala
1 5 10 15
Ala Pro Ser Gln Asn Pro Thr Arg Thr Gln Pro Leu Glu Lys Arg Ala
20 25 30
Thr Thr Leu Cys Gly Gln Trp Asp Ser Val Glu Thr Gly Gly Tyr Thr
35 40 45
Ile Tyr Asn Asn Leu Trp Gly Gln Asp Asn Gly Ser Gly Ser Gln Cys
50 55 60
Leu Thr Val Glu Gly Val Thr Asp Gly Leu Ala Ala Trp Ser Ser Thr
65 70 75 80
Trp Ser Trp Ser Gly Gly Ser Ser Ser Val Lys Ser Tyr Ser Asn Ala
85 90 95
Val Leu Ser Ala Glu Ala Ala Arg Ile Ser Ala Ile Ser Ser Ile Pro
100 105 110
Ser Lys Trp Glu Trp Ser Tyr Thr Gly Thr Asp Ile Val Ala Asn Val
115 120 125
Ala Tyr Asp Leu Phe Ser Asn Thr Asp Cys Gly Asp Thr Pro Glu Tyr
130 135 140
Glu Ile Met Ile Trp Leu Ser Ala Leu Gly Gly Ala Gly Pro Ile Ser
145 150 155 160
Ser Thr Gly Ser Ser Ile Ala Thr Val Thr Ile Ala Gly Ala Ser Trp
165 170 175
Asn Leu Trp Gln Gly Gln Asn Asn Gln Met Thr Val Phe Ser Phe Val
180 185 190
Ala Glu Ser Asp Gln Lys Ser Phe Ser Gly Asp Leu Asn Asp Phe Ile
195 200 205
Gln Tyr Leu Val Asp Ser Gln Gly Tyr Ser Gly Ser Gln Cys Leu Tyr
210 215 220
Ser Ile Gly Ala Gly Thr Glu Pro Phe Thr Gly Thr Asp Ala Glu Phe
225 230 235 240
Ile Thr Thr Gly Tyr Ser Val Ser Val Ser Ala Gly Asp Ser Gly Ser
245 250 255
Asp Glu Thr Thr Ser Ser Gln Ala Gln Ser Ser Thr Val Glu Thr
260 265 270
Ser Thr Ala Thr Gln Pro Gln Ser Ser Ser Thr Val Val Pro Thr Val
275 280 285

Thr Leu Ser Gln Pro Ser Asn Glu Ser Thr Thr Thr Pro Val Gln Ser
 290 295 300
 Gln Pro Ser Ser Val Glu Thr Thr Pro Thr Ala Gln Pro Gln Ser Ser
 305 310 315 320
 Ser Val Gln Thr Thr Thr Thr Ala Gln Ala Gln Pro Thr Pro Glu Arg
 325 330 335
 Ala Ala Pro Asp Ala Gly Ser Ala Glu Leu Leu Ser Ser Ala Thr Met
 340 345 350
 His Leu Asp Arg Arg
 355

<210> 24

<211> 247

<212> PRT

<213> *Emericella desertorum*

<400> 24

Met Lys Leu Leu Ala Leu Ser Leu Val Ser Leu Ala Ser Ala Ala Ser
 1 5 10 15
 Ala Ala Ser Ile Leu Ser Asn Thr Phe Thr Arg Arg Ser Asp Phe Cys
 20 25 30
 Gly Gln Trp Asp Thr Ala Thr Val Gly Asn Phe Ile Val Tyr Asn Asn
 35 40 45
 Leu Trp Gly Gln Asp Asn Ala Asp Ser Gly Ser Gln Cys Thr Gly Val
 50 55 60
 Asp Ser Ala Asn Gly Asn Ser Ile Ser Trp His Thr Thr Trp Ser Trp
 65 70 75 80
 Ser Gly Gly Ser Ser Ser Val Lys Ser Tyr Ala Asn Ala Ala Tyr Gln
 85 90 95
 Phe Thr Ser Thr Lys Leu Asn Ser Leu Ser Ser Ile Pro Thr Ser Trp
 100 105 110
 Lys Trp Gln Tyr Ser Thr Thr Asp Ile Val Ala Asn Val Ala Tyr Asp
 115 120 125
 Leu Phe Thr Ser Ser Ser Ala Gly Gly Asp Ser Glu Tyr Glu Ile Met
 130 135 140
 Ile Trp Leu Ala Ala Leu Gly Gly Ala Gly Pro Ile Ser Ser Thr Gly
 145 150 155 160
 Ser Ser Ile Ala Thr Val Thr Leu Gly Gly Val Thr Trp Ser Leu Tyr
 165 170 175
 Ser Gly Pro Asn Gly Ser Met Gln Val Tyr Ser Phe Val Ala Ser Ser
 180 185 190
 Thr Thr Glu Ser Phe Ser Ala Asp Leu Met Asp Phe Ile Asn Tyr Leu
 195 200 205
 Ala Glu Asn Gln Gly Leu Ser Ser Ser Gln Tyr Leu Thr His Val Gln
 210 215 220
 Ala Gly Thr Glu Pro Phe Thr Gly Thr Asp Ala Thr Leu Thr Val Ser
 225 230 235 240
 Ser Tyr Ser Val Ser Val Ser
 245

<210> 25

<211> 244

<212> PRT

<213> *Fusarium solani*

<400> 25

Met Lys Ser Ala Ile Val Ala Ala Leu Ala Gly Leu Ala Ala Ala Ser

1				5					10					15			
Pro	Thr	Arg	Leu	Ile	Pro	Arg	Gly	Gln	Phe	Cys	Gly	Gln	Trp	Asp	Ser		
			20					25					30				
Glu	Thr	Ala	Gly	Ala	Tyr	Thr	Ile	Tyr	Asn	Asn	Leu	Trp	Gly	Lys	Asp		
		35					40					45					
Asn	Ala	Glu	Ser	Gly	Glu	Gln	Cys	Thr	Thr	Asn	Ser	Gly	Glu	Gln	Ser		
	50					55					60						
Asp	Gly	Ser	Ile	Ala	Trp	Ser	Val	Glu	Trp	Ser	Trp	Thr	Gly	Gly	Gln		
65					70					75					80		
Gly	Gln	Val	Lys	Ser	Tyr	Pro	Asn	Ala	Val	Val	Glu	Ile	Glu	Lys	Lys		
			85					90					95				
Thr	Leu	Gly	Glu	Val	Ser	Ser	Ile	Pro	Ser	Ala	Trp	Asp	Trp	Thr	Tyr		
		100						105					110				
Thr	Gly	Asn	Gly	Ile	Ile	Ala	Asn	Val	Ala	Tyr	Asp	Leu	Phe	Thr	Ser		
		115					120					125					
Ser	Thr	Glu	Ser	Gly	Asp	Ala	Glu	Tyr	Glu	Phe	Met	Ile	Trp	Leu	Ser		
	130					135					140						
Ala	Leu	Gly	Gly	Ala	Gly	Pro	Ile	Ser	Asn	Asp	Gly	Ser	Pro	Val	Ala		
145					150					155					160		
Thr	Val	Glu	Leu	Ala	Gly	Thr	Ser	Trp	Lys	Leu	Tyr	Gln	Gly	Lys	Asn		
				165				170					175				
Asn	Gln	Met	Thr	Val	Phe	Ser	Phe	Val	Ala	Glu	Ser	Asp	Val	Asn	Asn		
			180					185					190				
Phe	Cys	Gly	Asp	Leu	Ala	Asp	Phe	Thr	Asp	Tyr	Leu	Val	Asp	Asn	His		
	195					200						205					
Gly	Val	Ser	Ser	Ser	Gln	Ile	Leu	Gln	Ser	Val	Gly	Ala	Gly	Thr	Glu		
	210				215						220						
Pro	Phe	Glu	Gly	Thr	Asn	Ala	Val	Phe	Thr	Thr	Asn	Asn	Tyr	His	Ala		
225					230					235					240		
Asp	Val	Glu	Tyr														

<210> 26

<211> 250

<212> .PRT

<213> Fusarium solani

<400> 26

Met	Lys	Phe	Phe	Gly	Val	Val	Ser	Ala	Phe	Leu	Ala	Ala	Thr	Ala	Val		
1				5					10					15			
Ala	Thr	Pro	Thr	Thr	Pro	Thr	Glu	Thr	Ile	Glu	Lys	Arg	Asp	Thr	Thr		
		20					25						30				
Trp	Cys	Asp	Ala	Phe	Gly	Ser	Leu	Ala	Thr	Ser	Gly	Tyr	Thr	Val	Tyr		
		35					40					45					
His	Asn	Asn	Trp	Gly	Lys	Gly	Asp	Ala	Thr	Ser	Gly	Ser	Gln	Cys	Thr		
	50					55					60						
Thr	Phe	Thr	Ser	Val	Ser	Asn	Asn	Asn	Phe	Val	Trp	Ser	Thr	Ser	Trp		
65					70					75					80		
Thr	Trp	Ala	Gly	Gly	Ala	Gly	Lys	Val	Lys	Ser	Tyr	Ser	Asn	Val	Ala		
			85					90					95				
Leu	Glu	Lys	Ile	Asn	Lys	Lys	Ile	Ser	Asp	Ile	Lys	Ser	Val	Ser	Thr		
		100						105					110				
Arg	Trp	Ile	Trp	Arg	Tyr	Thr	Gly	Thr	Lys	Met	Ile	Ala	Asn	Val	Ser		
		115					120					125					
Tyr	Asp	Leu	Trp	Phe	Ala	Pro	Thr	Ala	Ser	Ser	Asn	Asn	Ala	Tyr	Glu		
	130					135					140						
Ile	Met	Ile	Trp	Val	Gly	Ala	Tyr	Gly	Gly	Ala	Leu	Pro	Ile	Ser	Thr		
145					150					155					160		

Pro Gly Lys Gly Val Ile Asp Arg Pro Thr Leu Ala Gly Ile Pro Trp
 165 170 175
 Asp Val Tyr Lys Gly Pro Asn Gly Asp Val Thr Val Ile Ser Phe Val
 180 185 190
 Ala Ser Ser Asn Gln Gly Asn Phe Gln Ala Asp Leu Lys Glu Phe Leu
 195 200 205
 Asn Tyr Leu Thr Ser Lys Gln Gly Leu Pro Ser Asn Tyr Val Ala Thr
 210 215 220
 Ser Phe Gln Ala Gly Thr Glu Pro Phe Glu Gly Thr Asn Ala Val Leu
 225 230 235 240
 Lys Thr Ser Ala Tyr Thr Ile Ser Val Asn
 245 250

<210> 27

<211> 371

<212> PRT

<213> Streptomyces sp. 11AG8

<400> 27

Met Arg Ser His Pro Arg Ser Ala Thr Met Thr Val Leu Val Val Leu
 1 5 10 15
 Ala Ser Leu Gly Ala Leu Leu Thr Ala Ala Ala Pro Ala Gln Ala Asn
 20 25 30
 Gln Gln Ile Cys Asp Arg Tyr Gly Thr Thr Thr Ile Gln Asp Arg Tyr
 35 40 45
 Val Val Gln Asn Asn Arg Trp Gly Thr Ser Ala Thr Gln Cys Ile Asn
 50 55 60
 Val Thr Gly Asn Gly Phe Glu Ile Thr Gln Ala Asp Gly Ser Val Pro
 65 70 75 80
 Thr Asn Gly Ala Pro Lys Ser Tyr Pro Ser Val Tyr Asp Gly Cys His
 85 90 95
 Tyr Gly Asn Cys Ala Pro Arg Thr Thr Leu Pro Met Arg Ile Ser Ser
 100 105 110
 Ile Gly Ser Ala Pro Ser Ser Val Ser Tyr Arg Tyr Thr Gly Asn Gly
 115 120 125
 Val Tyr Asn Ala Ala Tyr Asp Ile Trp Leu Asp Pro Thr Pro Arg Thr
 130 135 140
 Asn Gly Val Asn Arg Thr Glu Ile Met Ile Trp Phe Asn Arg Val Gly
 145 150 155 160
 Pro Val Gln Pro Ile Gly Ser Pro Val Gly Thr Ala His Val Gly Gly
 165 170 175
 Arg Ser Trp Glu Val Trp Thr Gly Ser Asn Gly Ser Asn Asp Val Ile
 180 185 190
 Ser Phe Leu Ala Pro Ser Ala Ile Ser Ser Trp Ser Phe Asp Val Lys
 195 200 205
 Asp Phe Val Asp Gln Ala Val Ser His Gly Leu Ala Thr Pro Asp Trp
 210 215 220
 Tyr Leu Thr Ser Ile Gln Ala Gly Phe Glu Pro Trp Glu Gly Gly Thr
 225 230 235 240
 Gly Leu Ala Val Asn Ser Phe Ser Ser Ala Val Asn Ala Gly Gly Gly
 245 250 255
 Asn Gly Gly Thr Pro Gly Thr Pro Ala Ala Cys Gln Val Ser Tyr Ser
 260 265 270
 Thr His Thr Trp Pro Gly Gly Phe Thr Val Asp Thr Thr Ile Thr Asn
 275 280 285
 Thr Gly Ser Thr Pro Val Asp Gly Trp Glu Leu Asp Phe Thr Leu Pro
 290 295 300

Ala Gly His Thr Val Thr Ser Val Trp Asn Ala Leu Ile Ser Pro Ala
 305 310 315 320
 Ser Gly Ala Val Thr Ala Arg Ser Thr Gly Ser Asn Gly Arg Ile Ala
 325 330 335
 Ala Asn Gly Gly Thr Gln Ser Phe Gly Phe Gln Gly Thr Ser Ser Gly
 340 345 350
 Ala Gly Phe Thr Ala Pro Ala Gly Ala Arg Leu Asn Gly Thr Ser Cys
 355 360 365
 Thr Val Arg
 370

<210> 28
 <211> 221
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> consensus sequence

<220>
 <221> VARIANT
 <222> (1)...(221)
 <223> Xaa = Any Amino Acid

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 Cys Xaa Gln Tyr Xaa Xaa Xaa Xaa Xaa Xaa Gly Tyr Xaa Xaa Xaa Asn
 1 5 10 15
 Asn Xaa Trp Gly Xaa Xaa Xaa Xaa Xaa Ser Gly Xaa Gln Cys Thr Xaa
 20 25 30
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Trp Xaa Xaa Xaa Trp
 35 40 45
 Xaa Trp Ser Gly Gly Xaa Xaa Xaa Val Lys Ser Tyr Xaa Xaa Xaa Xaa
 50 55 60
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ser Xaa Ile Xaa Ser Xaa
 65 70 75 80
 Xaa Xaa Xaa Xaa Xaa Trp Xaa Tyr Xaa Gly Xaa Xaa Xaa Xaa Ala Asn
 85 90 95
 Val Ala Tyr Asp Leu Phe Thr Xaa Xaa Xaa Pro Xaa His Xaa Xaa Xaa
 100 105 110
 Xaa Gly Xaa Tyr Glu Xaa Met Ile Trp Leu Xaa Xaa Xaa Gly Gly Xaa
 115 120 125
 Xaa Pro Ile Gly Ser Xaa Xaa Xaa Xaa Val Xaa Xaa Xaa Xaa Xaa Xaa
 130 135 140
 Gly Xaa Xaa Trp Xaa Leu Xaa Xaa Gly Xaa Asn Gly Xaa Met Xaa Val
 145 150 155 160
 Xaa Ser Phe Val Ala Xaa Ser Ser Ser Ser Ser Ser Phe Xaa Gly Asp
 165 170 175
 Xaa Xaa Xaa Phe Xaa Xaa Tyr Leu Xaa Xaa Xaa Xaa Gly Xaa Pro Xaa
 180 185 190
 Xaa Xaa Gln Tyr Leu Xaa Xaa Xaa Gln Xaa Gly Thr Glu Pro Phe Thr
 195 200 205
 Gly Xaa Xaa Ala Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ala
 210 215 220

<210> 29
 <211> 25
 <212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 29

gaacgatggc aagggcggcg tgacg

25

<210> 30

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 30

cttctcgggc tgctacaacc caaacgg

27

<210> 31

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 31

acatcgtcga gtgttttggc acctac

26

<210> 32

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 32

catcgtcgag aactggggca cctacaacc

29

<210> 33

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 33

ggcacctacc gaccgtccac g

21

<210> 34

<211> 25

<212> DNA

<213> Artificial Sequence

<220>
 <223> synthetic oligonucleotide

 <400> 34
 caagctgggc gagcacacct ccgac 25

 <210> 35
 <211> 21
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> synthetic oligonucleotide

 <400> 35
 cgccgcaact gtcgctcgag c 21

 <210> 36
 <211> 29
 <212> DNA
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 <220>
 <223> synthetic oligonucleotide

 <400> 36
 gtggagggtt accaaagctc tggctctgc 29

 <210> 37
 <211> 27
 <212> DNA
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 <220>
 <223> synthetic oligonucleotide

 <400> 37
 tctggctctg cttgcatcac cgtcagc 27

 <210> 38
 <211> 27
 <212> DNA
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 <220>
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 <400> 38
 gagaagcgcc agtgcattca gcccggc 27

 <210> 39
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 <212> DNA
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 <220>

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<400> 39

gtgacgtact gcaatgggtcc cggcggg

27

<210> 40

<211> 33

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 40

ggcaccaaga acaggggtcat caacttctcg ggc

33

<210> 41

<211> 33

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 41

tccatcaccg tcagcgatta aagggggctc ttc

33

<210> 42

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 42

cccagacgat tcagtgcggc acgggctaca ac

32

<210> 43

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 43

cttctactcg tactggtgcg atggccacgg cg

32

<210> 44

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 44
 cgattcagcc cggtgcggc tacaacaacg gc 32

<210> 45
 <211> 35
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> synthetic oligonucleotide

<400> 45
 caacggctac ttctactgct actggaacga tggcc 35

<210> 46
 <211> 34
 <212> DNA
 <213> Artificial Sequence

<220>
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<400> 46
 ccggcacggg ctactgcaac ggctacttct actc 34

<210> 47
 <211> 31
 <212> DNA
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<220>
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<400> 47
 ggcgtgacgt acacctgcgg tcccggcggg c 31

<210> 48
 <211> 27
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> synthetic oligonucleotide

<400> 48
 ggcgccacca agtgcggcga ggtcacc 27

<210> 49
 <211> 28
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> synthetic oligonucleotide

<400> 49

gcgtgggctc agtgcggcct gacgctcg

28

<210> 50

<211> 752

<212> DNA

<213> *Trichoderma reesei*

<400> 50

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cccacaggcc	tgcagcctga	gagcagtgtc	aacgtcacag	agcgtggcat	gtacgacttt	120
gttcttggag	ctcacaatga	tcctcgccgt	cgtgctagca	tcaactacga	ccaaaactac	180
caaactggcg	gacaagtcag	ctattcgcc	tccaacactg	gcttctcagt	gaactggaac	240
actcaagatg	actttgttgt	gggcgttggt	tggacgactg	gatcttctgc	gtaggaggac	300
tcctcatcat	tctgcacttt	gaaagcatct	tctgaccaa	agcttctctt	agtcccatca	360
actttggcgg	ctcttttagt	gtcaacagcg	gaactggcct	gctttccgtc	tatggctgga	420
gcaccaaccc	actggttgag	tactacatca	tggaggacaa	ccacaactac	ccagcacagg	480
gtaccgtcaa	gggaaccgtc	accagcgacg	gagccactta	caccatctgg	gagaataccc	540
gtgtcaacga	gccttccatc	cagggcacag	cgaccttcaa	ccagtacatt	tccgtgcgga	600
actgcgccag	gaccagcgga	actgttactg	tgcagaacca	cttcaatgct	tgggcctcgc	660
ttggcctgca	ccttgggcag	atgaactacc	aggttgctgc	tgtcgaaggc	tggggtggtg	720
gtggttctgc	ctcacagagt	gtcagcaact	ag			752

<210> 51

<211> 248

<212> PRT

<213> *Trichoderma reesei*

<400> 51

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Thr	Leu	Ala	Met	Pro	Thr	Gly	Leu	Glu	Pro	Glu	Ser	Ser	Val	Asn
			20				25					30		Val
Thr	Glu	Arg	Gly	Met	Tyr	Asp	Phe	Val	Leu	Gly	Ala	His	Asn	Asp
		35					40					45		His
Arg	Arg	Arg	Ala	Ser	Ile	Asn	Tyr	Asp	Gln	Asn	Tyr	Gln	Thr	Gly
		50				55					60			Gly
Gln	Val	Ser	Tyr	Ser	Pro	Ser	Asn	Thr	Gly	Phe	Ser	Val	Asn	Trp
					70					75				Asn
Thr	Gln	Asp	Asp	Phe	Val	Val	Gly	Val	Gly	Trp	Thr	Thr	Gly	Ser
				85					90					Ser
Ala	Glu	Asp	Ser	Ser	Ser	Phe	Cys	Thr	Leu	Lys	Ala	Ser	Ser	Asp
			100					105					110	Gln
Lys	Leu	Leu	Leu	Val	Pro	Ser	Thr	Leu	Ala	Ala	Leu	Leu	Val	Ser
		115					120					125		Thr
Ala	Glu	Leu	Ala	Cys	Phe	Pro	Ser	Met	Ala	Gly	Ala	Pro	Thr	His
	130					135					140			Trp
Leu	Ser	Thr	Thr	Ser	Trp	Arg	Thr	Thr	Thr	Thr	Thr	Gln	His	Arg
	145				150					155				Val
Pro	Ser	Arg	Glu	Pro	Ser	Pro	Ala	Thr	Glu	Pro	Leu	Thr	Pro	Ser
			165						170				175	Gly
Arg	Ile	Pro	Val	Ser	Thr	Ser	Leu	Pro	Ser	Arg	Ala	Gln	Arg	Pro
			180				185						190	Ser
Thr	Ser	Thr	Phe	Pro	Cys	Gly	Thr	Arg	Pro	Gly	Pro	Ala	Glu	Leu
		195					200					205		Leu
Leu	Cys	Arg	Thr	Thr	Ser	Met	Leu	Gly	Pro	Arg	Leu	Ala	Cys	Thr
	210					215					220			Leu
Gly	Arg	Thr	Thr	Arg	Leu	Ser	Leu	Ser	Lys	Ala	Gly	Val	Val	Val

225
Leu Pro His Arg Val Ser Ala Thr
245

230

235

240